

Application No.: 09/671,773

**NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES**

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

- ☒ 1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.
- ☐ 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).
- ☐ 3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
- ☐ 4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
- ☐ 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
- ☐ 6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
- ☐ 7. Other: \_\_\_\_\_

**Applicant Must Provide:**

- ☒ An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
- ☒ An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
- ☒ A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).

For questions regarding compliance to these requirements, please contact:

For Rules Interpretation, call (703) 308-4216

For CRF Submission Help, call (703) 308-4212

For PatentIn software help, call (703) 308-6856

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## RAW SEQUENCE LISTING

DATE: 11/27/2002

PATENT APPLICATION: US/09/671,773A

TIME: 09:21:27

Input Set : A:\EP.txt

Output Set: N:\CRF4\11272002\I671773A.raw

3 <110> APPLICANT: BAYER CORPORATION  
 5 <120> TITLE OF INVENTION: Pituitary Adenylate Cyclase Activating Peptide (PACAP)  
 Receptor 3 (R3)  
 6 Agonists and Their Pharmacological Methods of Use  
 8 <130> FILE REFERENCE: Bayer  
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/671,773A  
 C--> 10 <141> CURRENT FILING DATE: 2001-05-31  
 10 <150> PRIOR APPLICATION NUMBER: 09/407,832  
 11 <151> PRIOR FILING DATE: 1999-09-28  
 13 <150> PRIOR APPLICATION NUMBER: 09/595,280  
 14 <151> PRIOR FILING DATE: 2000-06-15  
 16 <160> NUMBER OF SEQ ID NOS: 341  
 18 <170> SOFTWARE: PatentIn version 3.1  
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 22 <212> TYPE: PRT  
 23 <213> ORGANISM: Artificial Sequence  
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 26 <223> OTHER INFORMATION: synthetic construct  
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 29 <221> NAME/KEY: MOD\_RES  
 30 <222> LOCATION: (28)..(28)  
 31 <223> OTHER INFORMATION: AMIDATION  
 34 <400> SEQUENCE: 1  
 36 His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln  
 37 1 5 10 15  
 40 Met Ala Val Lys Lys Tyr Leu Asn Ser Ile Leu Asn  
 41 20 25  
 44 <210> SEQ ID NO: 2  
 45 <211> LENGTH: 38  
 46 <212> TYPE: PRT  
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 53 <221> NAME/KEY: MOD\_RES  
 54 <222> LOCATION: (38)..(38)  
 55 <223> OTHER INFORMATION: AMIDATION  
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 61 1 5 10 15  
 64 Met Ala Val Lys Lys Tyr Leu Ala Ala Val Leu Gly Lys Arg Tyr Lys  
 65 20 25 30  
 68 Gln Arg Val Lys Asn Lys

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83 <223> OTHER INFORMATION: AMIDATION
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88 His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly
89 1          5          10          15
92 Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg
93          20          25          30
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106 <222> LOCATION: (39)..(39)
107 <223> OTHER INFORMATION: AMIDATION
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116 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
117          20          25          30
120 Ser Gly Ala Pro Pro Pro Ser
121          35
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133 <221> NAME/KEY: MOD_RES
134 <222> LOCATION: (1)..(1)
135 <223> OTHER INFORMATION: ACETYLTATION
138 <220> FEATURE:
139 <221> NAME/KEY: MOD_RES
140 <222> LOCATION: (31)..(31)
141 <223> OTHER INFORMATION: AMIDATION
144 <220> FEATURE:
145 <221> NAME/KEY: MISC_FEATURE

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147 <223> OTHER INFORMATION: Nle
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152 His Ser Asp Ala Val Phe Thr Glu Asn Tyr Thr Lys Leu Arg Lys Gln
153 1 5 10 15
W--> 156 Xaa Ala Ala Lys Lys Tyr Leu Asn Asp Leu Lys Lys Gly Gly Thr
157 20 25 30
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176 <222> LOCATION: (1)..(1)
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182 His Ser Asp Ala Val Phe Thr Glu Asn Tyr Thr Lys Leu Arg Lys Gln
183 1 5 10 15
186 Leu Ala Ala Lys Lys Tyr Leu Asn Asp Leu Lys Lys Gly Gly Thr
187 20 25 30
190 <210> SEQ ID NO: 7
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192 <212> TYPE: PRT
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200 <222> LOCATION: (1)..(1)
201 <223> OTHER INFORMATION: ACETYLTATION
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206 His Ser Asp Ala Val Phe Thr Glu Asn Tyr Thr Lys Leu Arg Lys Gln
207 1 5 10 15
210 Leu Ala Ala Lys Lys Tyr Leu Asn Asp Leu Lys Lys Gly Gly Thr
211 20 25 30
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216 <212> TYPE: PRT
217 <213> ORGANISM: Artificial Sequence
219 <220> FEATURE:
220 <223> OTHER INFORMATION: synthetic construct
222 <220> FEATURE:
223 <221> NAME/KEY: PEPTIDE

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225 <223> OTHER INFORMATION:
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230 1 5 10 15
233 Leu Ala Ala Lys Lys Tyr Leu Asn Asp Leu Lys Lys Gly Gly Thr
234 20 25 30
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239 <212> TYPE: PRT
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248 <223> OTHER INFORMATION: Nle
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254 <223> OTHER INFORMATION: ACETYLTATION
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260 <223> OTHER INFORMATION: AMIDATION
263 <220> FEATURE:
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265 <222> LOCATION: (10)..(10)
266 <223> OTHER INFORMATION: CH3O-Y
269 <400> SEQUENCE: 9
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272 1 5 10 15
W--> 275 Xaa Ala Ala Lys Lys Tyr Leu Asn Asp Leu Lys Lys
276 20 25
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294 His Ser Asp Ala Val Phe Thr Glu Asn Tyr Thr Lys Leu Arg Lys Gln
295 1 5 10 15
298 Leu Ala Ala Lys Lys Tyr Leu Asn Asp Leu Lys Lys
299 20 25

```

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313 <223> OTHER INFORMATION: AMIDATION
316 <400> SEQUENCE: 11
318 His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln
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322 Met Ala Val Lys Lys Tyr Leu Asn Ser Ile Lys Lys
323          20          25
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327 <211> LENGTH: 31
328 <212> TYPE: PRT
329 <213> ORGANISM: Artificial Sequence
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336 <222> LOCATION: (1)..(31)
337 <223> OTHER INFORMATION:
W--> 339 <400> 12
341 His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln
342 1          5          10          15
345 Met Ala Val Lys Lys Tyr Leu Asn Ser Ile Lys Lys Gly Gly Thr
346          20          25          30
349 <210> SEQ ID NO: 13
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351 <212> TYPE: PRT
352 <213> ORGANISM: Artificial Sequence
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360 <223> OTHER INFORMATION:
W--> 362 <400> 13
364 His Ser Asp Ala Val Phe Thr Glu Asn Tyr Thr Lys Leu Arg Lys Gln
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368 Leu Ala Ala Lys Lys Tyr Leu Asn Asp Leu Leu Asn Gly Gly Thr
369          20          25          30
372 <210> SEQ ID NO: 14
373 <211> LENGTH: 31
374 <212> TYPE: PRT
375 <213> ORGANISM: Artificial Sequence

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RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/671,773A

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Input Set : A:\EP.txt  
Output Set: N:\CRF4\11272002\I671773A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; Xaa Pos. 17/  
Seq#:9; Xaa Pos. 10,17/

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Input Set : A:\EP.txt

Output Set: N:\CRF4\11272002\I671773A.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No  
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:156 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:16  
L:227 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:8,Line#:225  
L:271 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0  
L:275 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:16  
L:292 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:10,Line#:290  
L:339 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:12,Line#:337  
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L:385 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:14,Line#:383  
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L:551 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:21,Line#:549  
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L:689 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:27,Line#:687  
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Input Set : A:\EP.txt

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L:1306 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:54  
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